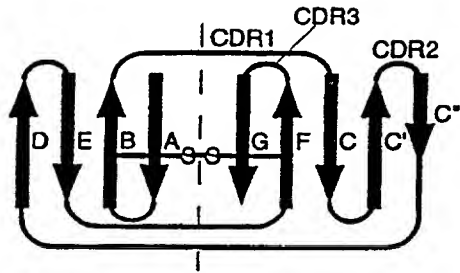


FIG. 1A



Immunoglobulin VH

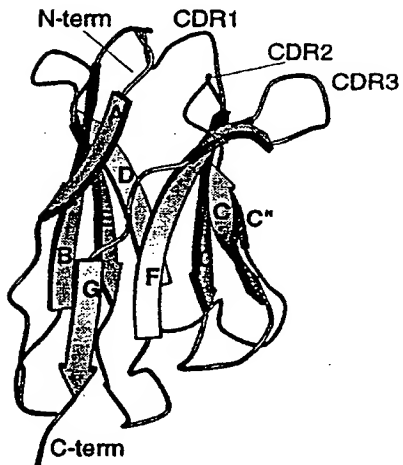
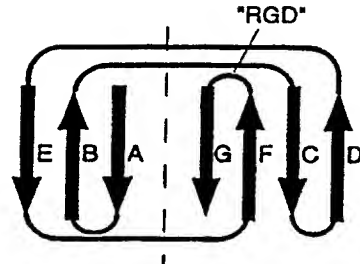


FIG. 1C

FIG. 1B



Fibronectin type III

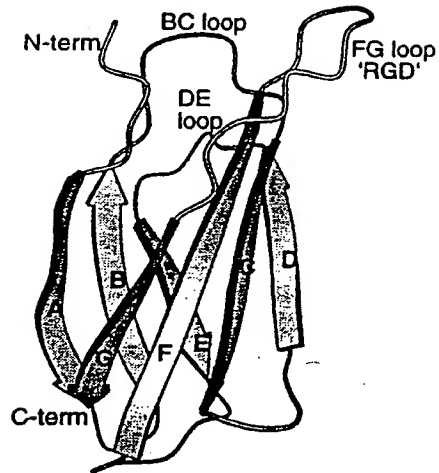


FIG. 1D

[illegible]

FIG. 2

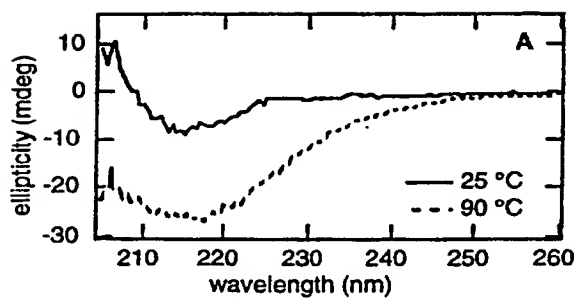


FIG. 3A

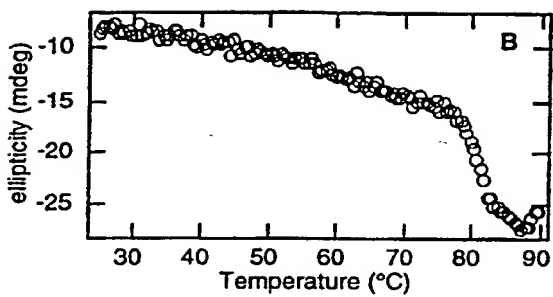


FIG. 3B

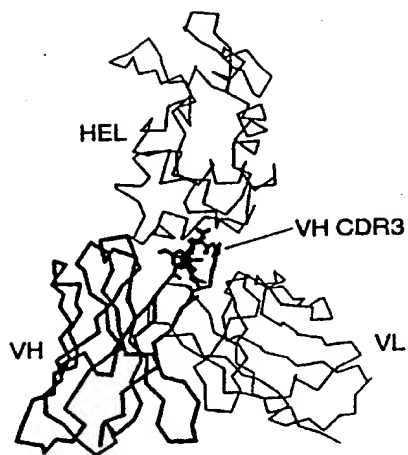


FIG. 4A

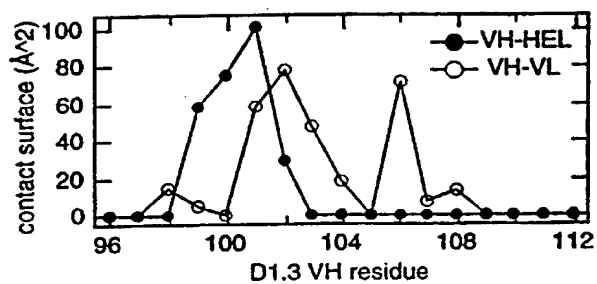


FIG. 4B

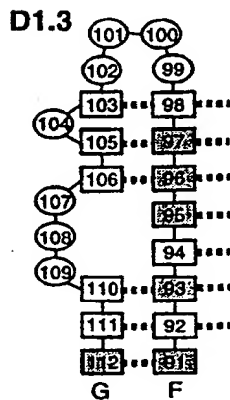


FIG. 4C

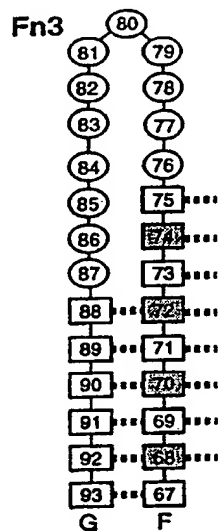


FIG. 4D

NdeI  
 CATATGCAGGTTTCTGATGTTCCGCGTGACCTGGAAGTTGTTGCTGCGACCCCGACTAGC  
 MetGlnValSerAspValProArgAspLeuGluValValAlaAlaThrProThrSer  
 -2 -1 1 10

BclI PvuII PstI BsiWI  
 CTGCTGATCAGCTGGGATGCTCCTGCAGTTACCGTGCGTTATTACCGTATCACGTACGGT  
 LeuLeuIleSerTrpAspAlaProAlaValThrValArgTyrTyrArgIleThrTyrGly  
 20 30

EcoRI  
 GAAACCGGTGGTAACTCCCCGGTTCAGGAATTCACCTGTACCTGGTTCCAAGTCTACTGCT  
 GluThrGlyGlyAsnSerProValGlnGluPheThrValProGlySerLysSerThrAla  
 40 50

SalI Bst1107I  
 ACCATCAGCGGCCTGAAACCGGGTGTGCGACTATACCATCACTGTATACGCTGTTACTGGC  
 ThrIleSerGlyLeuLysProGlyValAspTyrThrIleThrValTyrAlaValThrGly  
 60 70

SacI XhoI  
 CGTGGTGACAGCCCAGCGAGCTCCAAGCCAATCTCGATTAACCTACCGTACCTAGTAACTC  
 ArgGlyAspSerProAlaSerSerLysProIleSerIleAsnTyrArgThr  
 80 90

FIG. 5

090343-02404

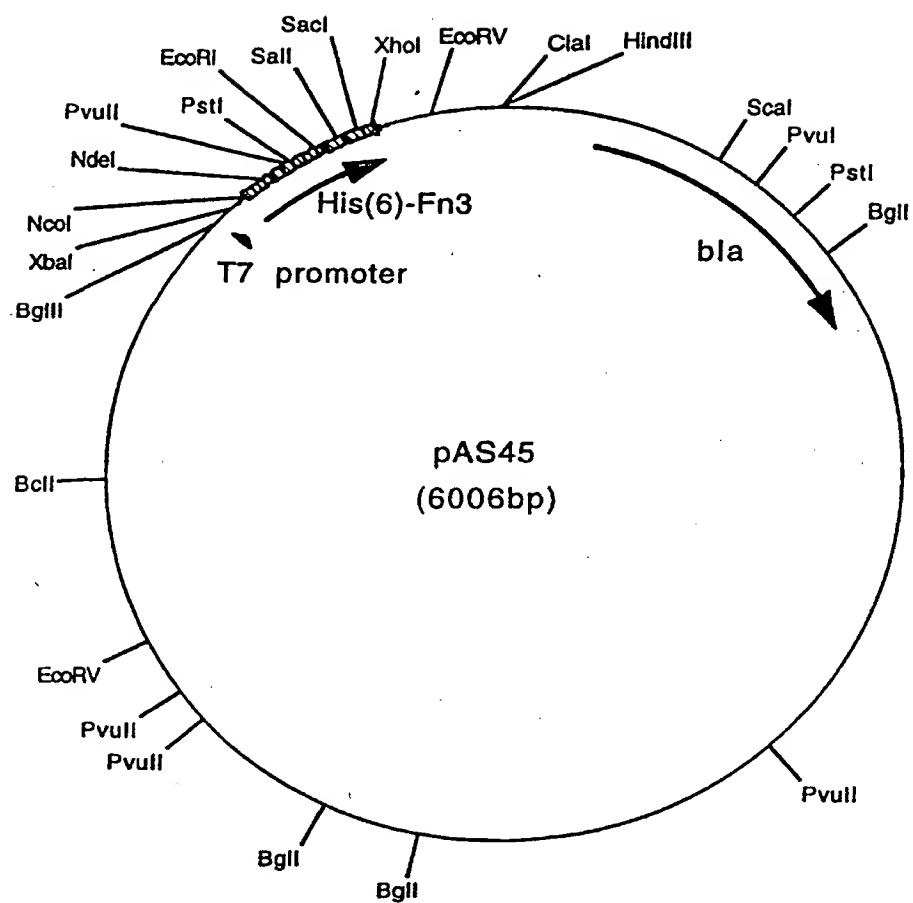


FIG. 6

A circular map of the pAS25 (5946bp) plasmid. The map shows various restriction enzyme sites around the circle. Key features include the T7 promoter (indicated by a small arrow), the Fn3 gene (indicated by a large arrow), and the bla gene (indicated by a large arrow). Restriction sites are labeled with names like EcoRI, SalI, SacI, XhoI, HindIII, ClaI, EcoRV, PstI, BglII, BclII, and PvuII.

FIG. 7

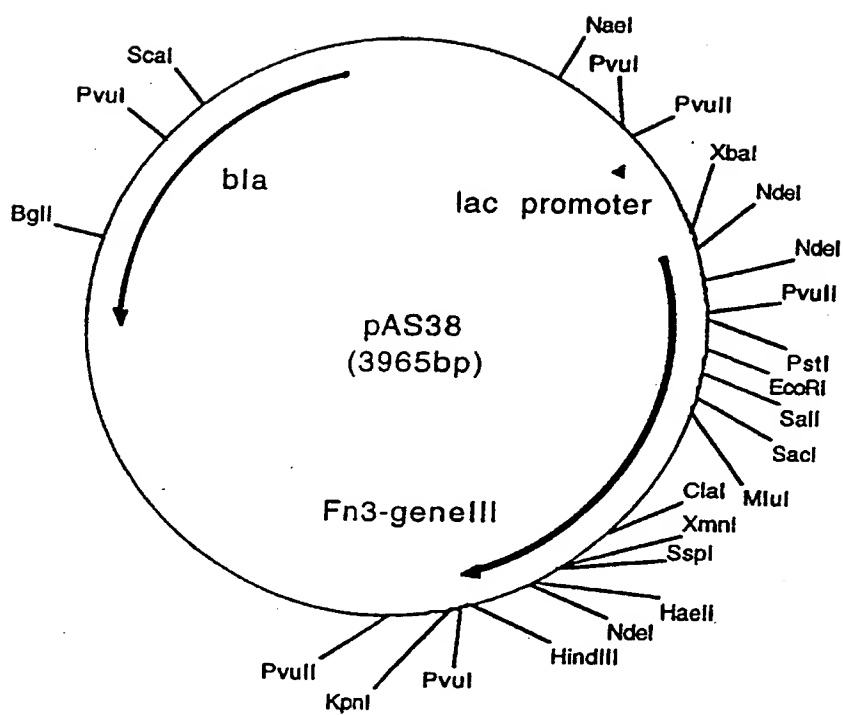


FIG. 8



phage ELISA (ubiquitin)

Clone #	Ligand (+)	Control
2-1	0.17	0.05
2-2	0.24	0.05
4-1	0.23	0.05
4-2	0.36	0.10

FIG. 9

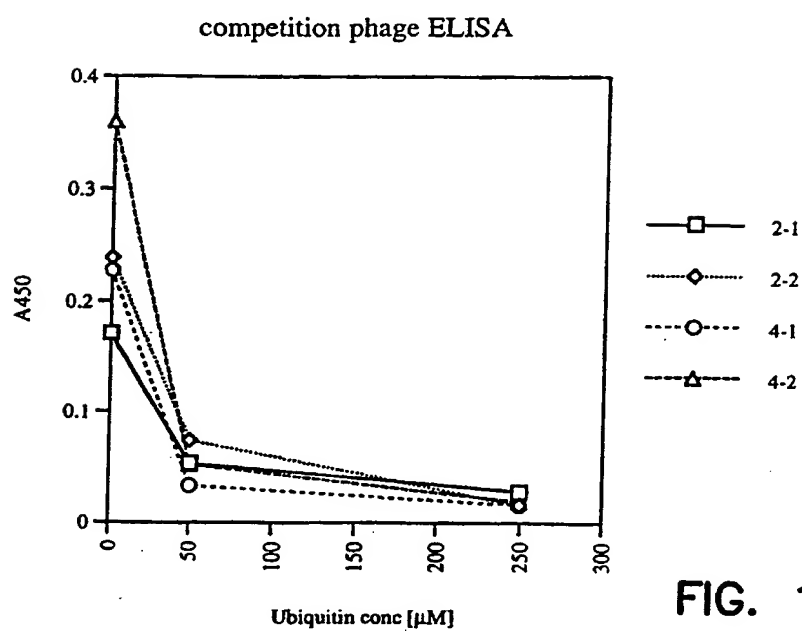


FIG. 10

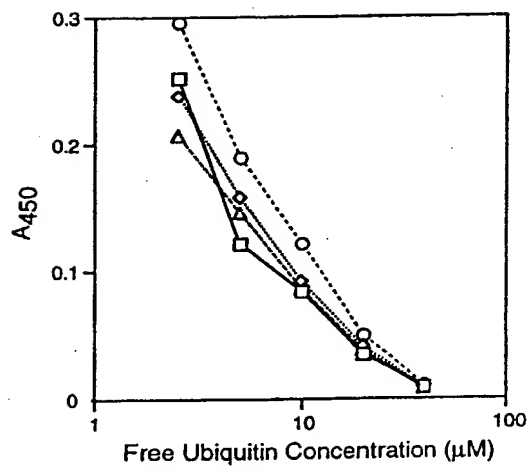
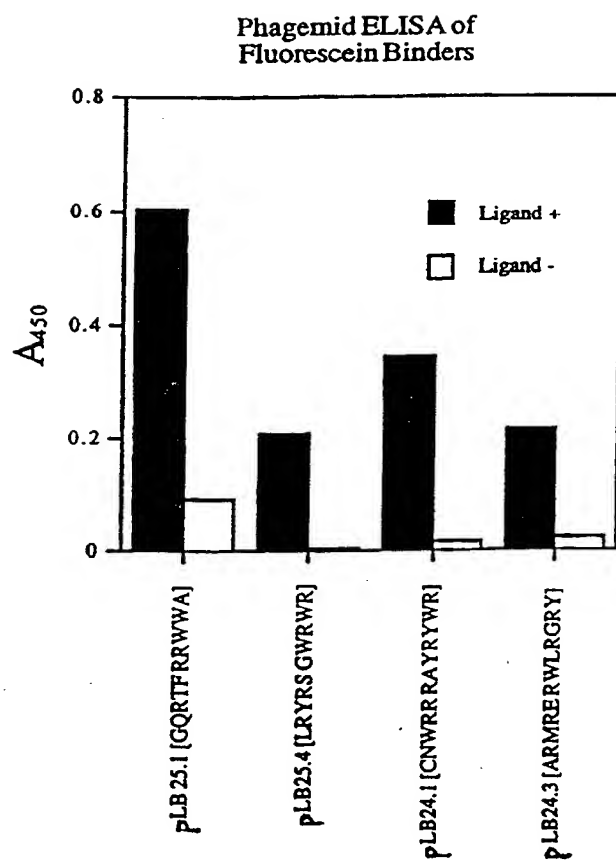
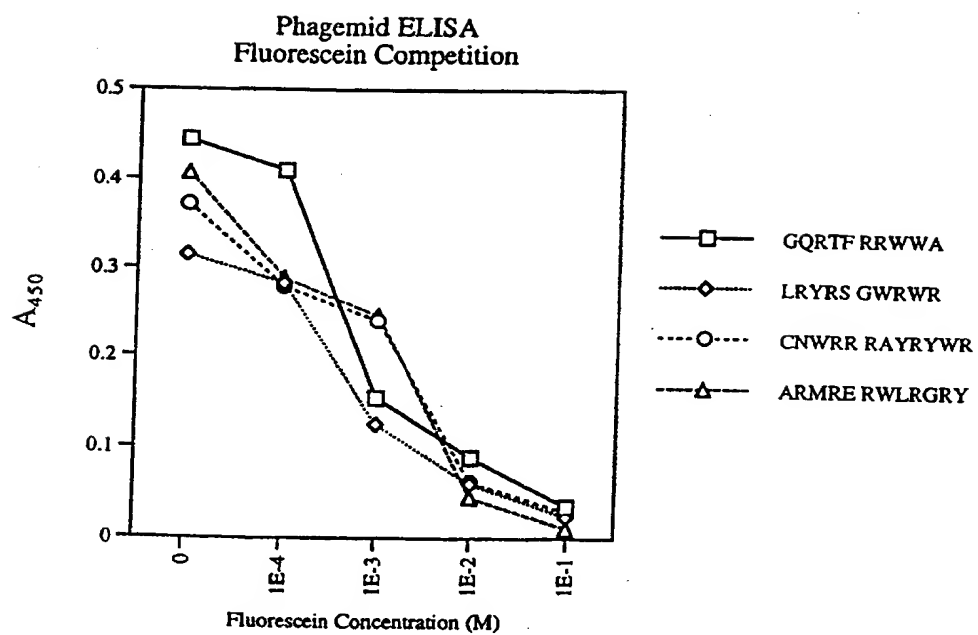


FIG. 11



**FIG. 12**



**FIG. 13**

FOFFCQ" ZHED0660

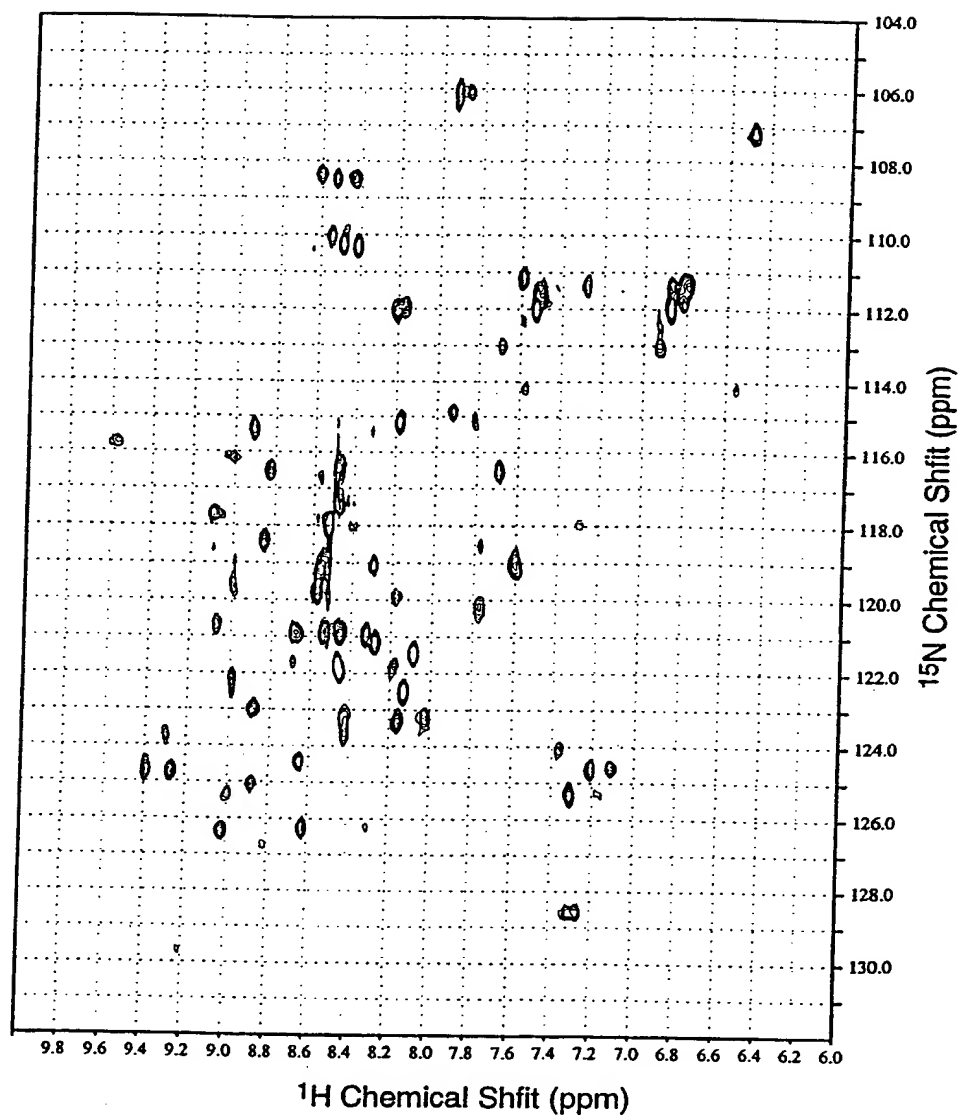


FIG. 14

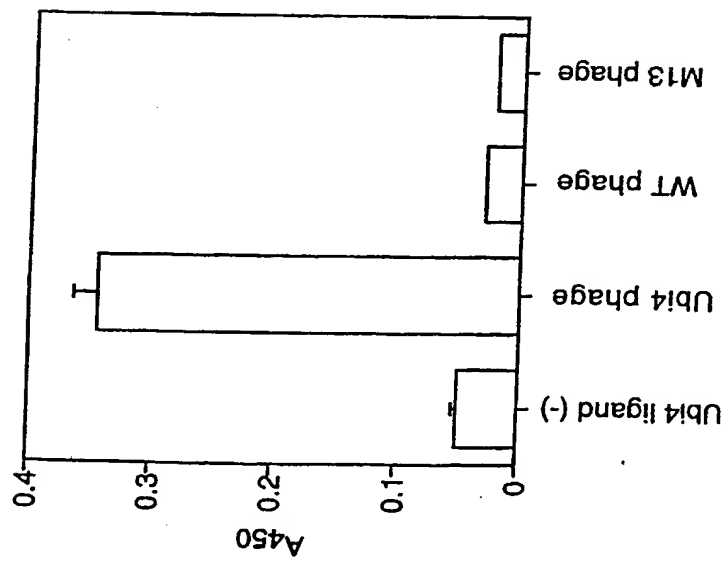


FIG. 15A

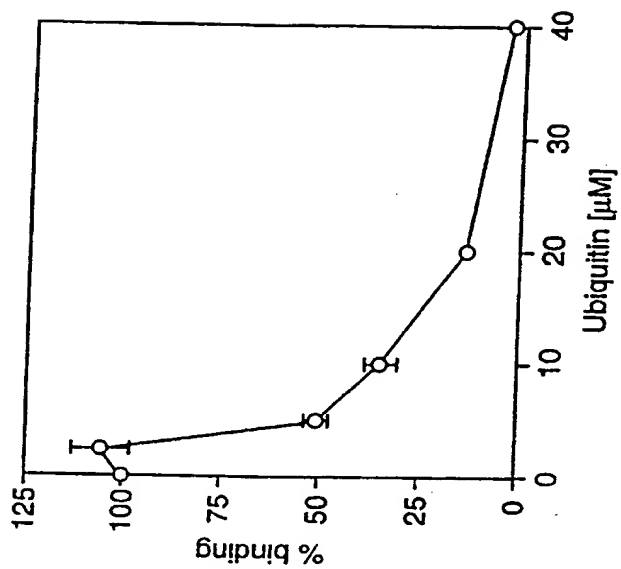


FIG. 15B

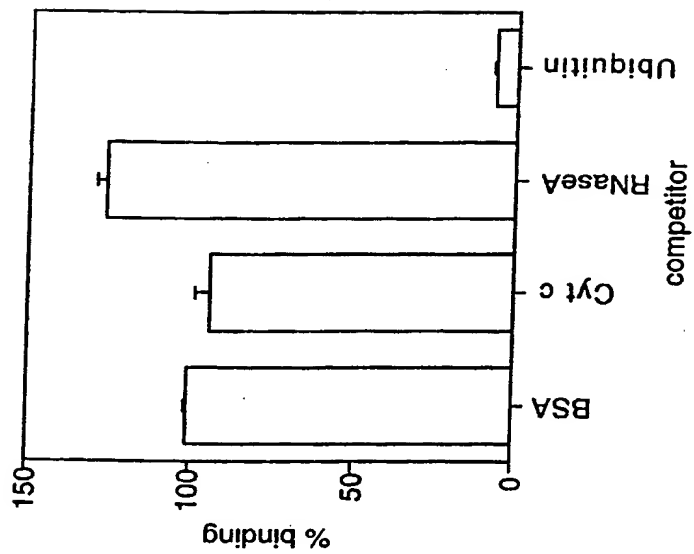


FIG. 15C

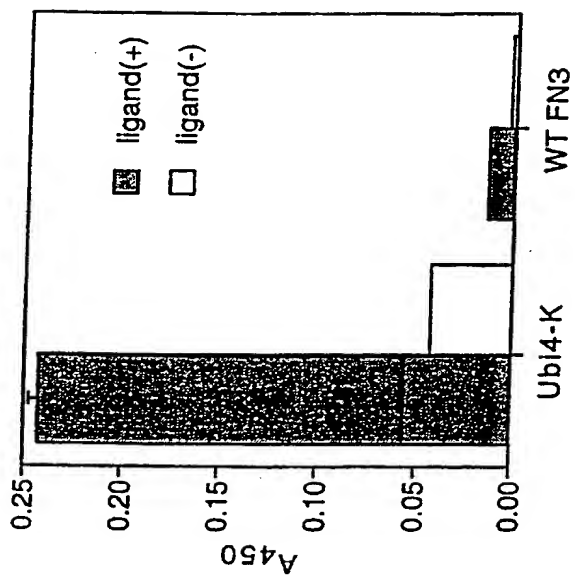


FIG. 15D

09903413 01101

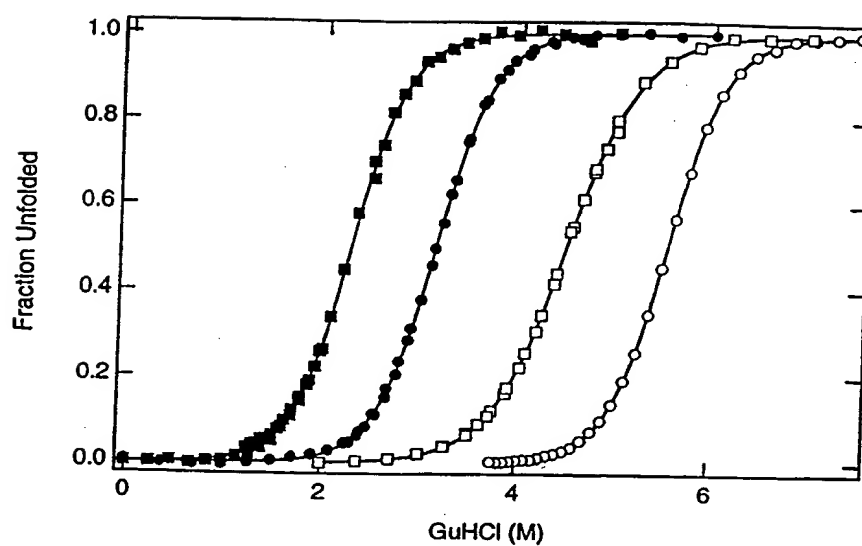


FIG. 16



A scatter plot showing the relationship between  $^1\text{H}$  and  $^{15}\text{N}$  chemical shifts for protein 1A28. The x-axis is labeled ' $^1\text{H}$  Chemical Shift (ppm)' and ranges from 10 to 6. The y-axis is labeled ' $^{15}\text{N}$  Chemical Shift (ppm)' and ranges from 110 to 130. The plot contains two sets of data points: open circles representing one set of assignments and solid dots representing another. The points are distributed across the plot, with a higher density in the central region (8-9 ppm  $^1\text{H}$ , 118-122 ppm  $^{15}\text{N}$ ).

FIG. 17A

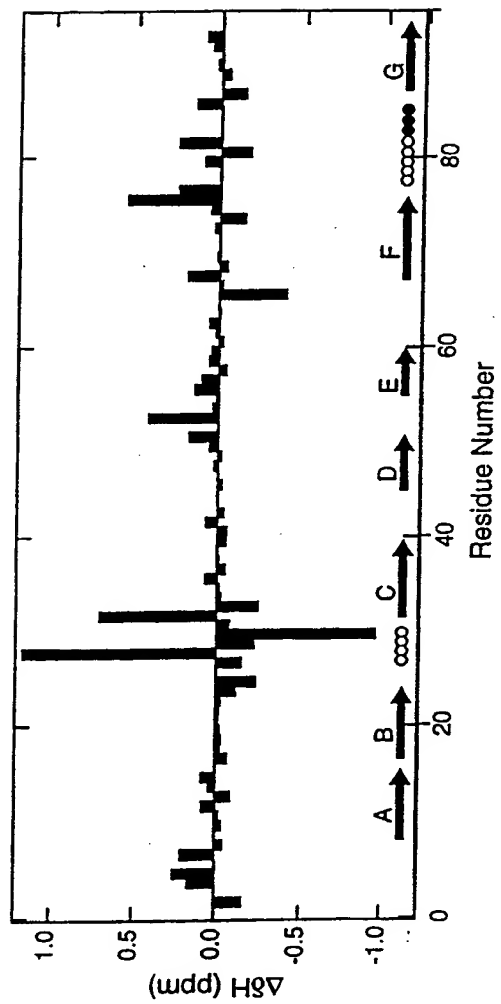


FIG. 17B

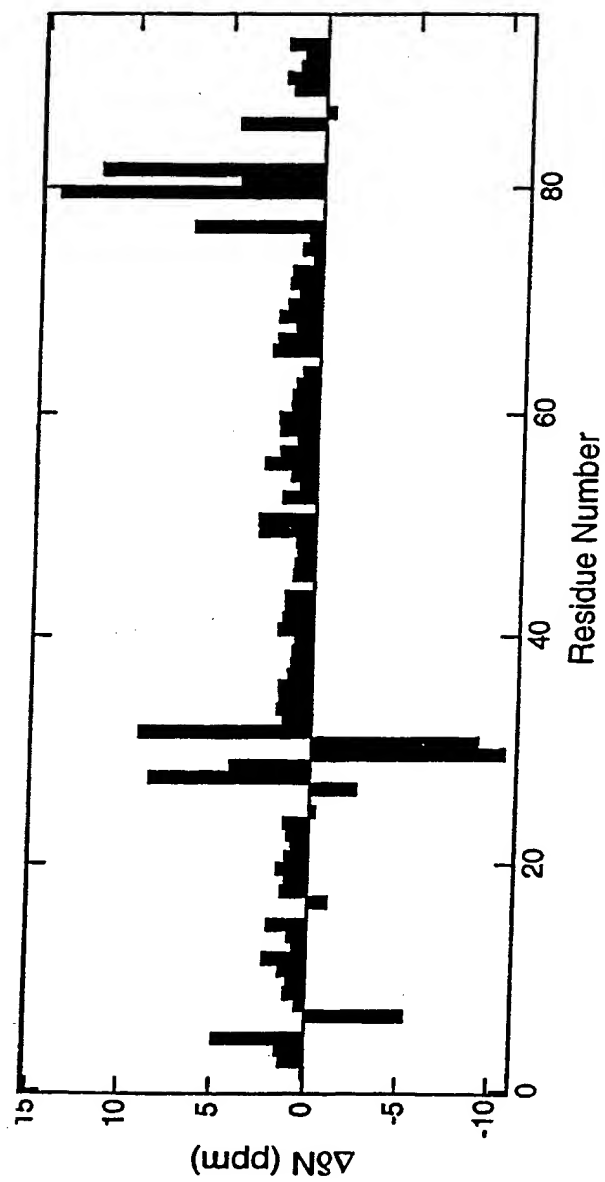


FIG. 17C

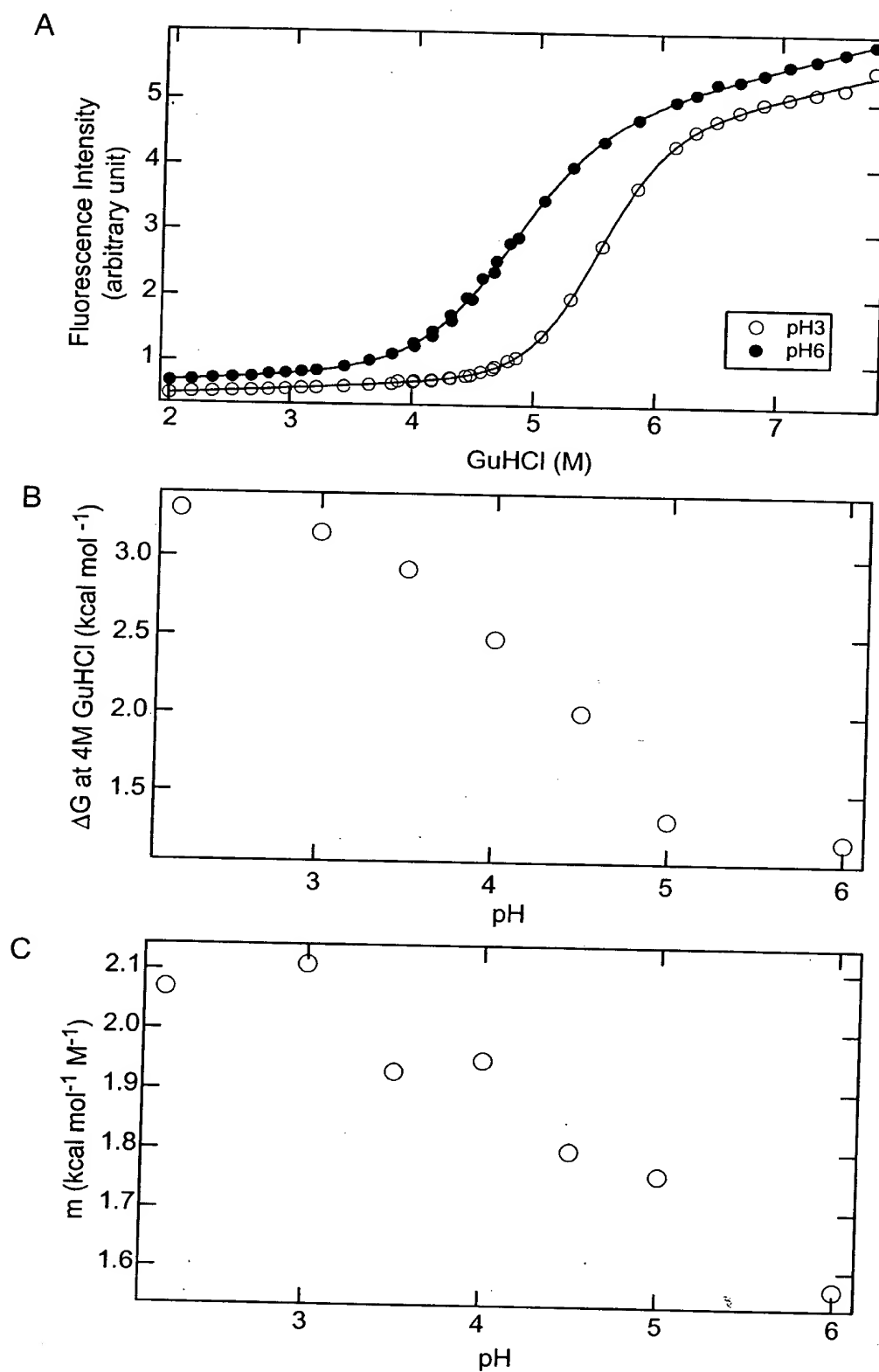


FIG. 18

00004400 at E0660

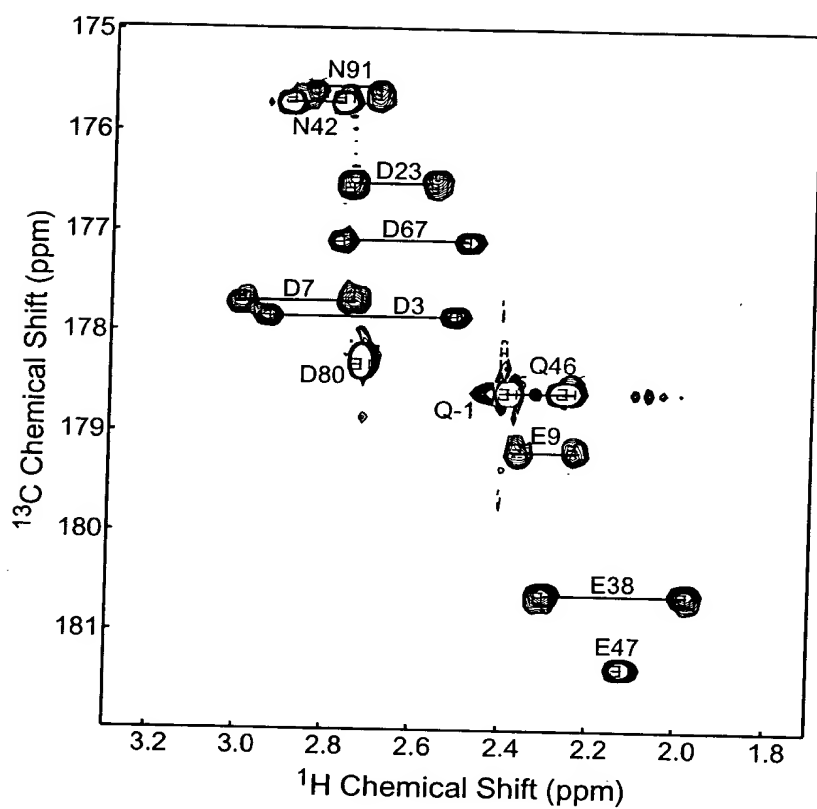


FIG. 19

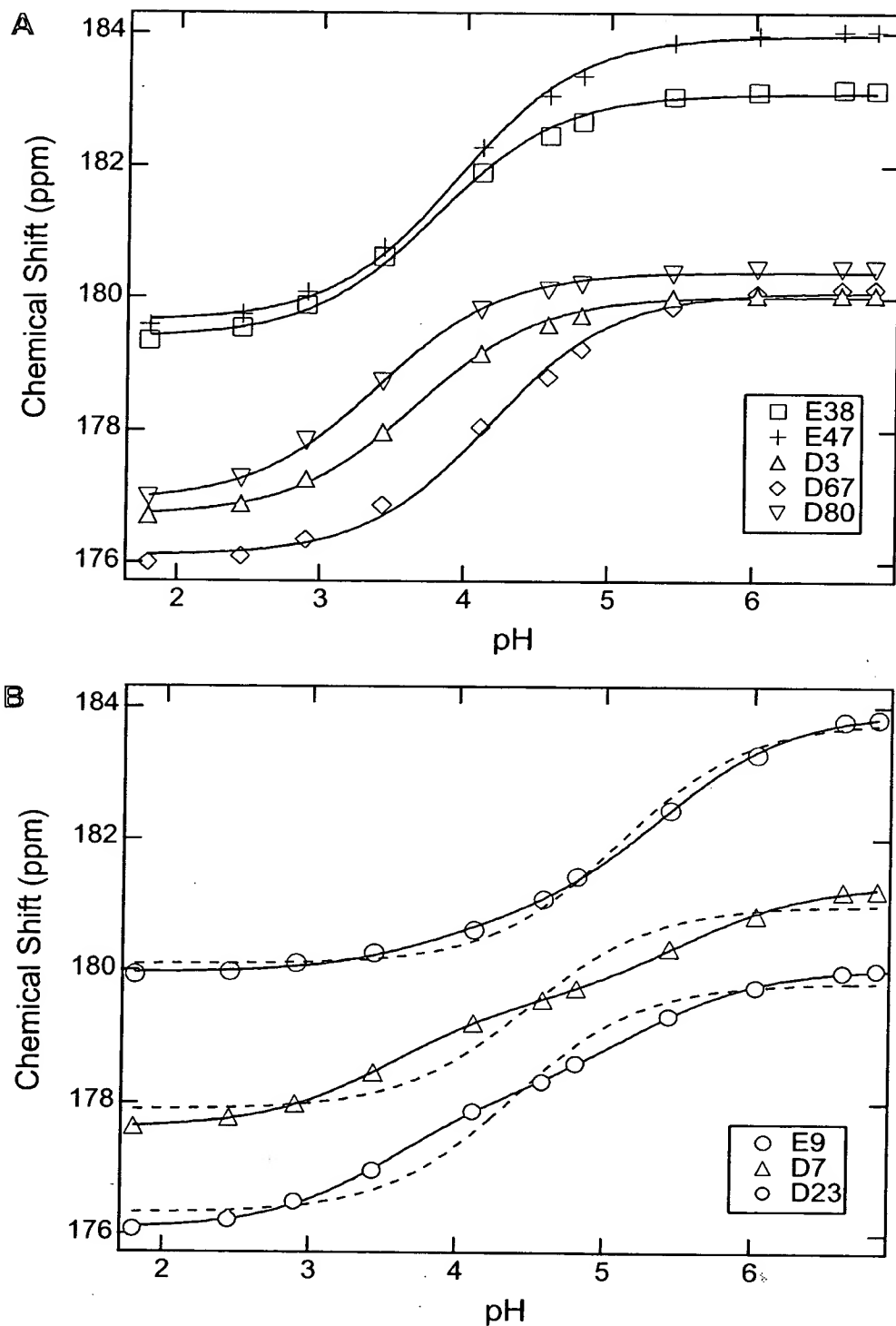


FIG. 20

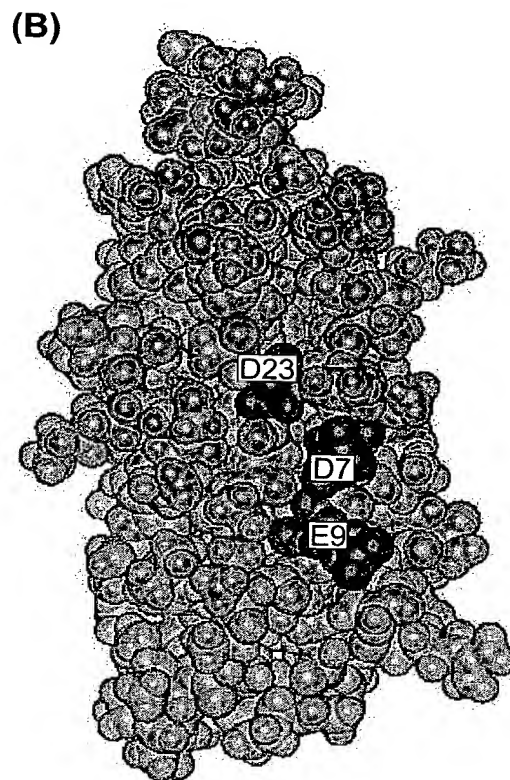
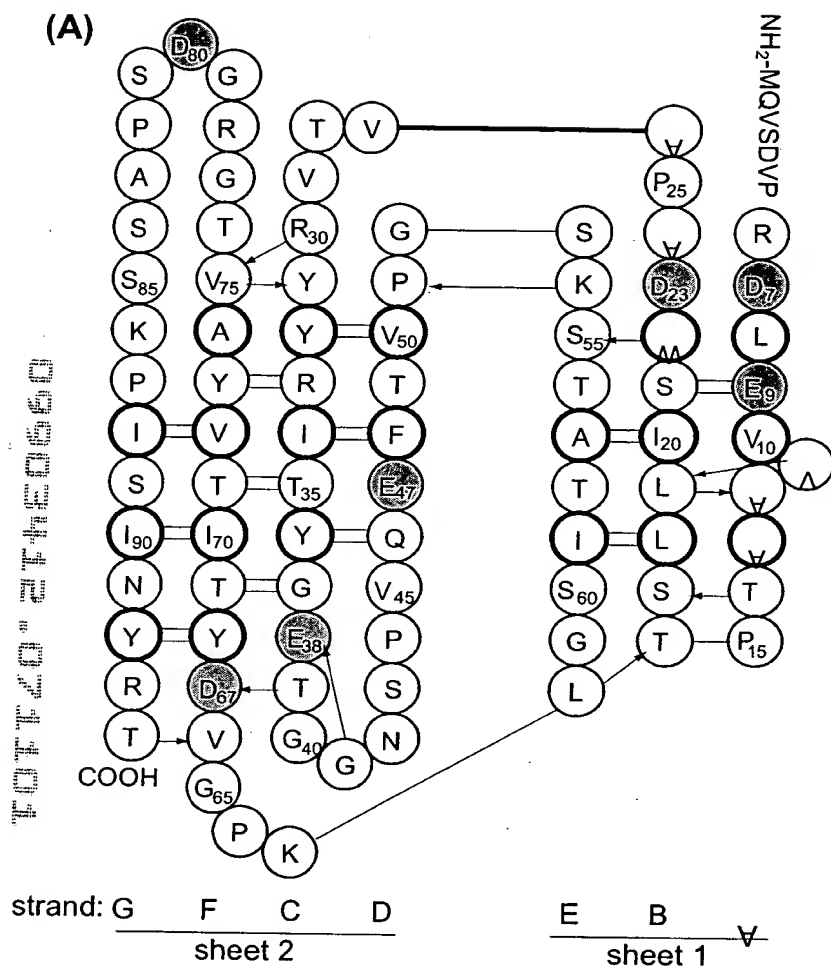


FIG. 21

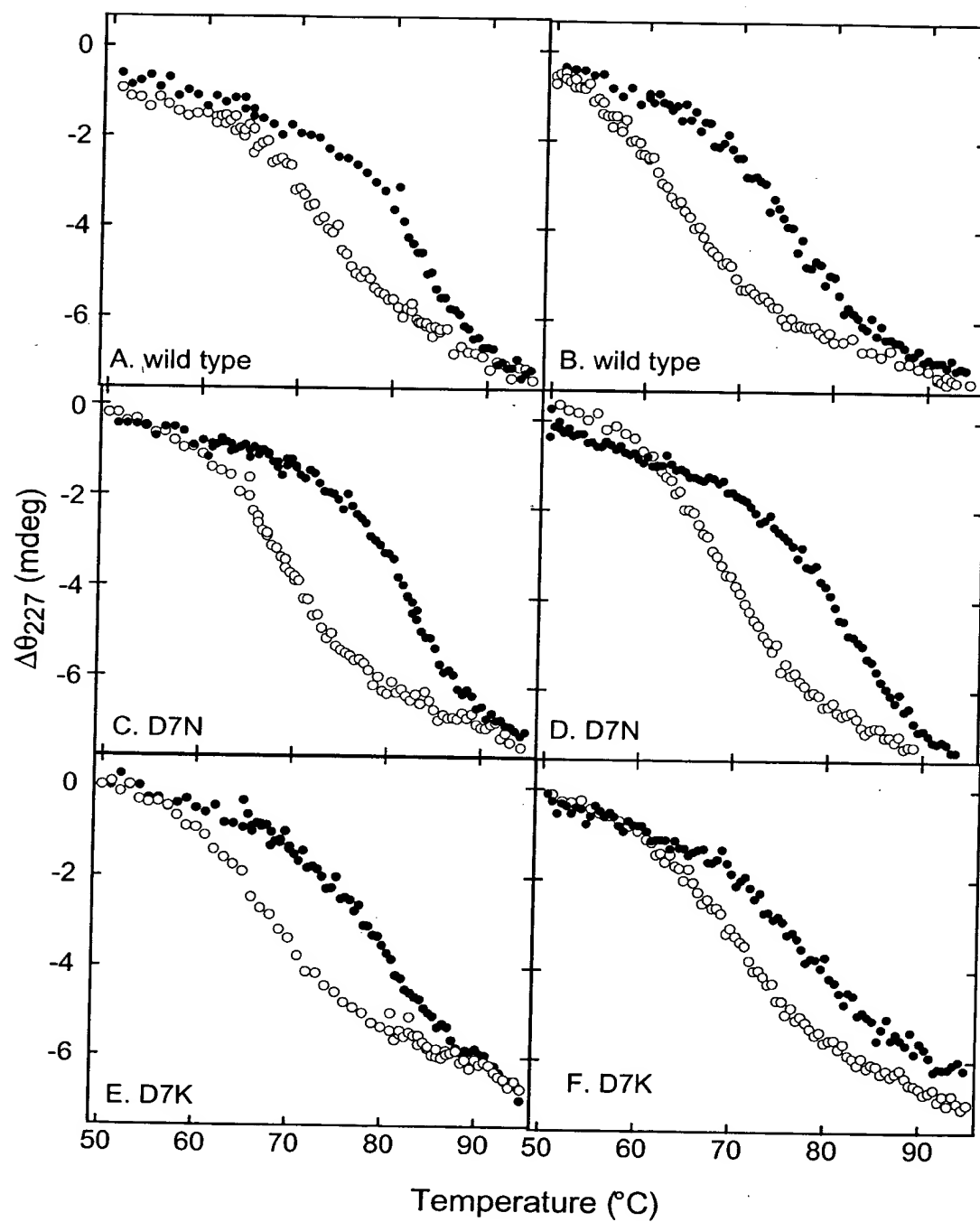


FIG. 22



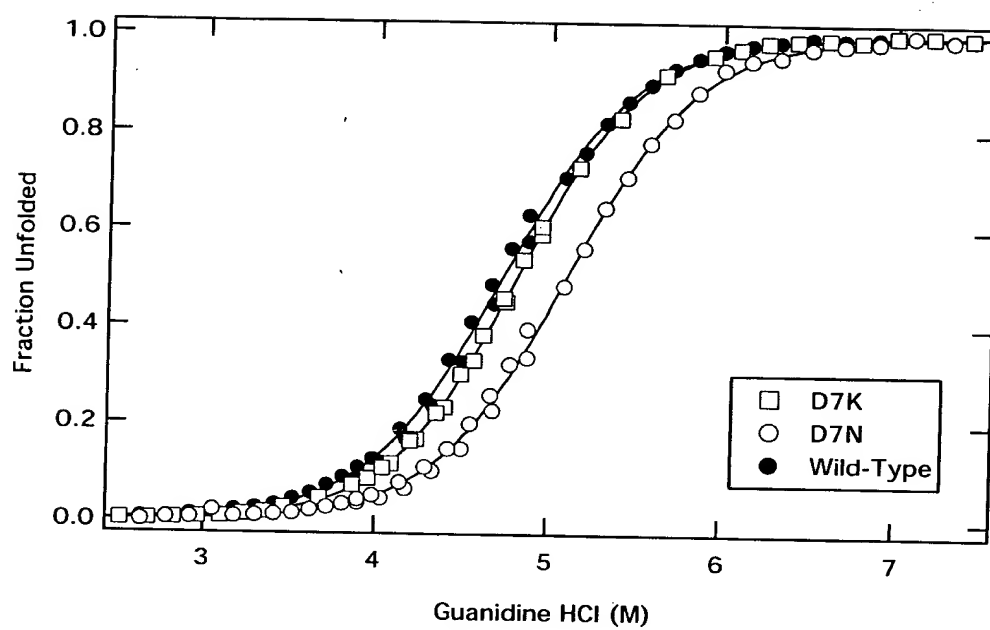


FIG. 23

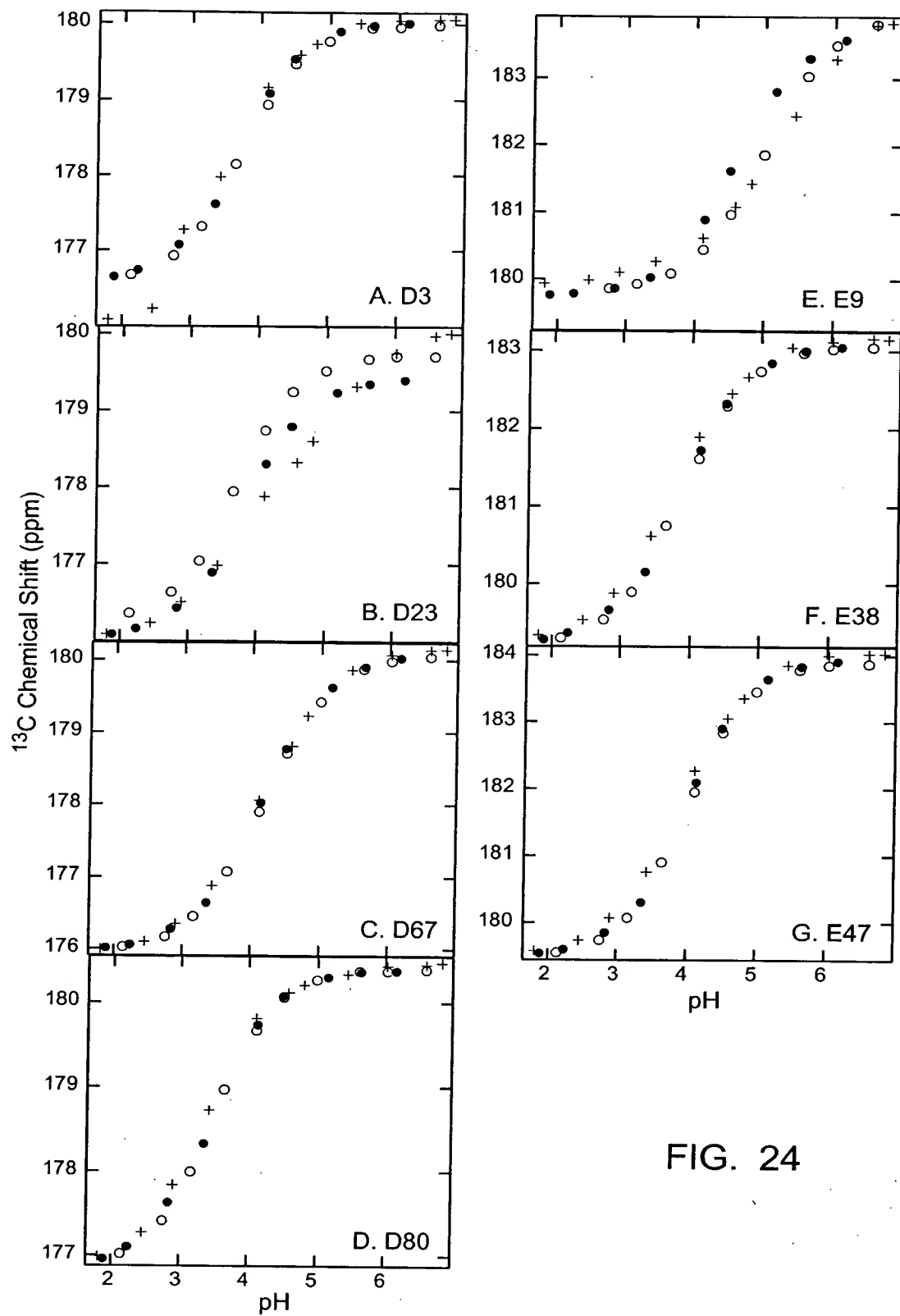


FIG. 24